



caa acg ccc tgg tct tgg gta acc gca tca cta cga aat ctt caa cct	433																																																						
Gln Thr Pro Trp Ser Trp Val Thr Ala Ser Leu Arg Asn Leu Gln Pro																																																							
130	135	140		gat aca gaa gct aac att tat aac tgg caa att aga cca cta tcc aac	481	Asp Thr Glu Ala Asn Ile Tyr Asn Trp Gln Ile Arg Pro Leu Ser Asn		145	150	155		att gcg aac tgg caa aac cta aaa gtt gga tgt gct cac aaa gtg tgc	529	Ile Ala Asn Trp Gln Asn Leu Lys Val Gly Cys Ala His Lys Val Cys		160	165	170		aaa ttc ccc acc ggg aca aat atg gtt gtg tct tgc gct tat ggc ggc	577	Lys Phe Pro Thr Gly Thr Asn Met Val Val Ser Cys Ala Tyr Gly Gly		175	180	185		gaa gta ctc caa gat aac gaa gtt gta tgg gac aag gga cca act tgc	625	Glu Val Leu Gln Asp Asn Glu Val Val Trp Asp Lys Gly Pro Thr Cys		190	195	200	205	atg tgc aat gct tat ccc aac tcg ttc tgc tgc aac aat ctg tgt gac	673	Met Cys Asn Ala Tyr Pro Asn Ser Phe Cys Cys Asn Asn Leu Cys Asp		210	215	220		aca ata gct gct gcg aca ctt cgc aag cag cct tgt aaa tcg act tga	721	Thr Ile Ala Ala Thr Leu Arg Lys Gln Pro Cys Lys Ser Thr		225	230	235		agcgaaaagg cgttggtgat gtcccgaaaga gaacggaagt gatcacatca cagtatccca	781	taatgtcggtt catcataata aacgcacttc tctgaaaaaaaaaaaaaaaa	828
140																																																							
gat aca gaa gct aac att tat aac tgg caa att aga cca cta tcc aac	481																																																						
Asp Thr Glu Ala Asn Ile Tyr Asn Trp Gln Ile Arg Pro Leu Ser Asn																																																							
145	150	155		att gcg aac tgg caa aac cta aaa gtt gga tgt gct cac aaa gtg tgc	529	Ile Ala Asn Trp Gln Asn Leu Lys Val Gly Cys Ala His Lys Val Cys		160	165	170		aaa ttc ccc acc ggg aca aat atg gtt gtg tct tgc gct tat ggc ggc	577	Lys Phe Pro Thr Gly Thr Asn Met Val Val Ser Cys Ala Tyr Gly Gly		175	180	185		gaa gta ctc caa gat aac gaa gtt gta tgg gac aag gga cca act tgc	625	Glu Val Leu Gln Asp Asn Glu Val Val Trp Asp Lys Gly Pro Thr Cys		190	195	200	205	atg tgc aat gct tat ccc aac tcg ttc tgc tgc aac aat ctg tgt gac	673	Met Cys Asn Ala Tyr Pro Asn Ser Phe Cys Cys Asn Asn Leu Cys Asp		210	215	220		aca ata gct gct gcg aca ctt cgc aag cag cct tgt aaa tcg act tga	721	Thr Ile Ala Ala Thr Leu Arg Lys Gln Pro Cys Lys Ser Thr		225	230	235		agcgaaaagg cgttggtgat gtcccgaaaga gaacggaagt gatcacatca cagtatccca	781	taatgtcggtt catcataata aacgcacttc tctgaaaaaaaaaaaaaaaa	828								
155																																																							
att gcg aac tgg caa aac cta aaa gtt gga tgt gct cac aaa gtg tgc	529																																																						
Ile Ala Asn Trp Gln Asn Leu Lys Val Gly Cys Ala His Lys Val Cys																																																							
160	165	170		aaa ttc ccc acc ggg aca aat atg gtt gtg tct tgc gct tat ggc ggc	577	Lys Phe Pro Thr Gly Thr Asn Met Val Val Ser Cys Ala Tyr Gly Gly		175	180	185		gaa gta ctc caa gat aac gaa gtt gta tgg gac aag gga cca act tgc	625	Glu Val Leu Gln Asp Asn Glu Val Val Trp Asp Lys Gly Pro Thr Cys		190	195	200	205	atg tgc aat gct tat ccc aac tcg ttc tgc tgc aac aat ctg tgt gac	673	Met Cys Asn Ala Tyr Pro Asn Ser Phe Cys Cys Asn Asn Leu Cys Asp		210	215	220		aca ata gct gct gcg aca ctt cgc aag cag cct tgt aaa tcg act tga	721	Thr Ile Ala Ala Thr Leu Arg Lys Gln Pro Cys Lys Ser Thr		225	230	235		agcgaaaagg cgttggtgat gtcccgaaaga gaacggaagt gatcacatca cagtatccca	781	taatgtcggtt catcataata aacgcacttc tctgaaaaaaaaaaaaaaaa	828																
170																																																							
aaa ttc ccc acc ggg aca aat atg gtt gtg tct tgc gct tat ggc ggc	577																																																						
Lys Phe Pro Thr Gly Thr Asn Met Val Val Ser Cys Ala Tyr Gly Gly																																																							
175	180	185		gaa gta ctc caa gat aac gaa gtt gta tgg gac aag gga cca act tgc	625	Glu Val Leu Gln Asp Asn Glu Val Val Trp Asp Lys Gly Pro Thr Cys		190	195	200	205	atg tgc aat gct tat ccc aac tcg ttc tgc tgc aac aat ctg tgt gac	673	Met Cys Asn Ala Tyr Pro Asn Ser Phe Cys Cys Asn Asn Leu Cys Asp		210	215	220		aca ata gct gct gcg aca ctt cgc aag cag cct tgt aaa tcg act tga	721	Thr Ile Ala Ala Thr Leu Arg Lys Gln Pro Cys Lys Ser Thr		225	230	235		agcgaaaagg cgttggtgat gtcccgaaaga gaacggaagt gatcacatca cagtatccca	781	taatgtcggtt catcataata aacgcacttc tctgaaaaaaaaaaaaaaaa	828																								
185																																																							
gaa gta ctc caa gat aac gaa gtt gta tgg gac aag gga cca act tgc	625																																																						
Glu Val Leu Gln Asp Asn Glu Val Val Trp Asp Lys Gly Pro Thr Cys																																																							
190	195	200	205	atg tgc aat gct tat ccc aac tcg ttc tgc tgc aac aat ctg tgt gac	673	Met Cys Asn Ala Tyr Pro Asn Ser Phe Cys Cys Asn Asn Leu Cys Asp		210	215	220		aca ata gct gct gcg aca ctt cgc aag cag cct tgt aaa tcg act tga	721	Thr Ile Ala Ala Thr Leu Arg Lys Gln Pro Cys Lys Ser Thr		225	230	235		agcgaaaagg cgttggtgat gtcccgaaaga gaacggaagt gatcacatca cagtatccca	781	taatgtcggtt catcataata aacgcacttc tctgaaaaaaaaaaaaaaaa	828																																
200	205																																																						
atg tgc aat gct tat ccc aac tcg ttc tgc tgc aac aat ctg tgt gac	673																																																						
Met Cys Asn Ala Tyr Pro Asn Ser Phe Cys Cys Asn Asn Leu Cys Asp																																																							
210	215	220		aca ata gct gct gcg aca ctt cgc aag cag cct tgt aaa tcg act tga	721	Thr Ile Ala Ala Thr Leu Arg Lys Gln Pro Cys Lys Ser Thr		225	230	235		agcgaaaagg cgttggtgat gtcccgaaaga gaacggaagt gatcacatca cagtatccca	781	taatgtcggtt catcataata aacgcacttc tctgaaaaaaaaaaaaaaaa	828																																								
220																																																							
aca ata gct gct gcg aca ctt cgc aag cag cct tgt aaa tcg act tga	721																																																						
Thr Ile Ala Ala Thr Leu Arg Lys Gln Pro Cys Lys Ser Thr																																																							
225	230	235		agcgaaaagg cgttggtgat gtcccgaaaga gaacggaagt gatcacatca cagtatccca	781	taatgtcggtt catcataata aacgcacttc tctgaaaaaaaaaaaaaaaa	828																																																
235																																																							
agcgaaaagg cgttggtgat gtcccgaaaga gaacggaagt gatcacatca cagtatccca	781																																																						
taatgtcggtt catcataata aacgcacttc tctgaaaaaaaaaaaaaaaa	828																																																						

<210> 2  
<211> 236  
<212> PRT  
<213> *Ostertagia ostertagi*

<220>  
<221> misc\_feature  
<222> (37)..(37)  
<223> The 'Xaa' at location 37 stands for Lys, or Gln.

<220>  
<221> misc\_feature  
<222> (121)..(121)  
<223> The 'Xaa' at location 121 stands for Asn, or Thr.

<400> 2

Met Gln Ala Leu Ile Gly Ile Ala Ala Leu Tyr Leu Val Leu Val Thr  
1 5 10 15

Ser Asn Thr Glu Ala Gly Phe Cys Cys Pro Ala Asp Leu Asn Gln Thr  
20 25 30

Asp Glu Ala Arg Xaa Ile Phe Leu Asp Phe His Asn Gln Val Arg Arg  
35 40 45

Asp Ile Ala Gly Ala Ser Pro Leu Leu Asn Leu Thr Gly Ala Val Gln  
50 55 60

Met Arg Asn Val Leu Gly Pro Ala Lys Asn Met Tyr Arg Met Asp Trp  
65 70 75 80

Asp Cys Asn Leu Glu Ala Lys Ala Lys Met Ile Trp Pro Cys Thr  
85 90 95

Thr Pro Leu Pro Ile Asp Thr Ser Ile Pro Gln Asn Leu Ala Gln Trp  
100 105 110

Leu Leu Phe Gln Asn Ser Gln Glu Xaa Glu Val Leu Thr Gln Thr Pro  
115 120 125

Trp Ser Trp Val Thr Ala Ser Leu Arg Asn Leu Gln Pro Asp Thr Glu  
130 135 140

Ala Asn Ile Tyr Asn Trp Gln Ile Arg Pro Leu Ser Asn Ile Ala Asn  
145 150 155 160

Trp Gln Asn Leu Lys Val Gly Cys Ala His Lys Val Cys Lys Phe Pro  
165 170 175

Thr Gly Thr Asn Met Val Val Ser Cys Ala Tyr Gly Gly Glu Val Leu  
180 185 190

Gln Asp Asn Glu Val Val Trp Asp Lys Gly Pro Thr Cys Met Cys Asn  
195 200 205

Ala Tyr Pro Asn Ser Phe Cys Cys Asn Asn Leu Cys Asp Thr Ile Ala  
210 215 220

Ala Ala Thr Leu Arg Lys Gln Pro Cys Lys Ser Thr  
225 230 235

<210> 3  
<211> 306  
<212> DNA  
<213> Ostertagia ostertagi

<220>  
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<222> (3)...(284)

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<220>
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<222> (10)..(10)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (16)..(16)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (82)..(82)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (226)..(226)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (240)..(240)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (248)..(248)
<223> n is a, c, g, or t

<400> 3
gc ggc cgc gnc gac cnt gtg atc agc atc atg gct ctg tgg ccc gtg      47
  Gly Arg Xaa Asp Xaa Val Ile Ser Ile Met Ala Leu Trp Pro Val
  1           5           10          15
                                         15

gac cgt ttc gag cgc atg ctg gaa gag ccg ttc ang cgt gtg gat cgt      95
  Asp Arg Phe Glu Arg Met Leu Glu Pro Phe Xaa Arg Val Asp Arg
  20          25          30
                                         30

ttc tgc ccg atg aga gat gcg gac tgg atg agc cgt cag att atg ccc      143
  Phe Cys Pro Met Arg Asp Ala Asp Trp Met Ser Arg Gln Ile Met Pro
  35          40          45
                                         45

tac tgg aga gat gcc gat cac tct gtg ctt cat gtg gga aat caa aca      191
  Tyr Trp Arg Asp Ala Asp His Ser Val Leu His Val Gly Asn Gln Thr
  50          55          60
                                         60

aag gat gtc gtg aat gac gag aag aaa ttc gca gnc gct ttg gat gtg      239
  Lys Asp Val Val Asn Asp Glu Lys Lys Phe Ala Xaa Ala Leu Asp Val
  65          70          75
                                         75

nca cac ttn agg cca gaa gag ttg aag gta caa ttg gaa gtg acg      284
  Xaa His Xaa Arg Pro Glu Glu Leu Lys Val Gln Leu Glu Val Thr
  80          85          90
                                         90

tgaccttaca atcgaaggac at
                                         306

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<210> 4  
<211> 94  
<212> PRT  
<213> Ostertagia ostertagi

<220>  
<221> misc\_feature  
<222> (3)..(3)  
<223> The 'Xaa' at location 3 stands for Asp, Gly, Ala, or Val.

<220>  
<221> misc\_feature  
<222> (5)..(5)  
<223> The 'Xaa' at location 5 stands for His, Arg, Pro, or Leu.

<220>  
<221> misc\_feature  
<222> (27)..(27)  
<223> The 'Xaa' at location 27 stands for Lys, Arg, Thr, or Met.

<220>  
<221> misc\_feature  
<222> (75)..(75)  
<223> The 'Xaa' at location 75 stands for Asp, Gly, Ala, or Val.

<220>  
<221> misc\_feature  
<222> (80)..(80)  
<223> The 'Xaa' at location 80 stands for Thr, Ala, Pro, or Ser.

<220>  
<221> misc\_feature  
<222> (82)..(82)  
<223> The 'Xaa' at location 82 stands for Leu, or Phe.

<400> 4

Gly Arg Xaa Asp Xaa Val Ile Ser Ile Met Ala Leu Trp Pro Val Asp  
1 5 10 15

Arg Phe Glu Arg Met Leu Glu Glu Pro Phe Xaa Arg Val Asp Arg Phe  
20 25 30

Cys Pro Met Arg Asp Ala Asp Trp Met Ser Arg Gln Ile Met Pro Tyr  
35 40 45

Trp Arg Asp Ala Asp His Ser Val Leu His Val Gly Asn Gln Thr Lys  
50 55 60

Asp Val Val Asn Asp Glu Lys Lys Phe Ala Xaa Ala Leu Asp Val Xaa  
65 70 75 80

His Xaa Arg Pro Glu Glu Leu Lys Val Gln Leu Glu Val Thr  
85 90

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<210> 5
<211> 583
<212> DNA
<213> Ostertagia ostertagi
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<220>  
<221> CDS  
<222> (2) . . (583)

<400> 5  
 g gct ttt atc gga aaa ccc gca ccc gac ttc gcc aca aag gcc gtc tat 49  
 Ala Phe Ile Gly Lys Pro Ala Pro Asp Phe Ala Thr Lys Ala Val Tyr  
 1 5 10 15  
  
 aat ggc gac ttc atc gac gtg aaa ctg tct gac tac aag ggc aag tac 97  
 Asn Gly Asp Phe Ile Asp Val Lys Leu Ser Asp Tyr Lys Gly Lys Tyr  
 20 25 30  
  
 acc gtc ctc ttc ttc tat cca ctg gat ttc acg ttt gtc tgt cct acg 145  
 Thr Val Leu Phe Phe Tyr Pro Leu Asp Phe Thr Phe Val Cys Pro Thr  
 35 40 45  
  
 gaa atc atc gcc ttt tcc gac cgt gtc gaa gaa ttc aaa aaa atc gat 193  
 Glu Ile Ile Ala Phe Ser Asp Arg Val Glu Glu Phe Lys Lys Ile Asp  
 50 55 60  
  
 gct gcg gtc ctc gct tgt tca amt gat tcc gtt ttc tct cat ctg gcg 241  
 Ala Ala Val Leu Ala Cys Ser Xaa Asp Ser Val Phe Ser His Leu Ala  
 65 70 75 80  
  
 tgg atc aat act cct cgc aag atg ggc ggc ctt ggt gac atg aac att 289  
 Trp Ile Asn Thr Pro Arg Lys Met Gly Gly Leu Gly Asp Met Asn Ile  
 85 90 95  
  
 ccc gtt ctt gct gac acc aac cac caa att gca aag gac tat ggt gta 337  
 Pro Val Leu Ala Asp Thr Asn His Gln Ile Ala Lys Asp Tyr Gly Val  
 100 105 110  
  
 ctg aaa gaa gac gaa gga atc gct tac aga ggt ctt ttc att att gac 385  
 Leu Lys Glu Asp Glu Gly Ile Ala Tyr Arg Gly Leu Phe Ile Ile Asp  
 115 120 125  
  
 cct aag gga att ctg cga cag atc act gtc aat gac ctt cct gtc ggt 433  
 Pro Lys Gly Ile Leu Arg Gln Ile Thr Val Asn Asp Leu Pro Val Gly  
 130 135 140  
  
 cgc tct gtg gat gag act ctc cgt ctg gtg cag gcc ttc caa tac gtt 481  
 Arg Ser Val Asp Glu Thr Leu Arg Leu Val Gln Ala Phe Gln Tyr Val  
 145 150 155 160  
  
 gac aag cat ggt gag gtg tgc cca gct ggt tgg act cct gga aaa gct 529  
 Asp Lys His Gly Glu Val Cys Pro Ala Gly Trp Thr Pro Gly Lys Ala  
 165 170 175

acc atc aag cca ggt gtc aag gac agc aag gag tac ttc agc aaa gca 577  
 Thr Ile Lys Pro Gly Val Lys Asp Ser Lys Glu Tyr Phe Ser Lys Ala  
 180 185 190

aac taa 583  
 Asn

<210> 6  
 <211> 193  
 <212> PRT  
 <213> *Ostertagia ostertagi*

<220>  
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 <222> (72)...(72)  
 <223> The 'Xaa' at location 72 stands for Asn, or Thr.

<400> 6

Ala Phe Ile Gly Lys Pro Ala Pro Asp Phe Ala Thr Lys Ala Val Tyr  
 1 5 10 15

Asn Gly Asp Phe Ile Asp Val Lys Leu Ser Asp Tyr Lys Gly Lys Tyr  
 20 25 30

Thr Val Leu Phe Phe Tyr Pro Leu Asp Phe Thr Phe Val Cys Pro Thr  
 35 40 45

Glu Ile Ile Ala Phe Ser Asp Arg Val Glu Glu Phe Lys Lys Ile Asp  
 50 55 60

Ala Ala Val Leu Ala Cys Ser Xaa Asp Ser Val Phe Ser His Leu Ala  
 65 70 75 80

Trp Ile Asn Thr Pro Arg Lys Met Gly Gly Leu Gly Asp Met Asn Ile  
 85 90 95

Pro Val Leu Ala Asp Thr Asn His Gln Ile Ala Lys Asp Tyr Gly Val  
 100 105 110

Leu Lys Glu Asp Glu Gly Ile Ala Tyr Arg Gly Leu Phe Ile Ile Asp  
 115 120 125

Pro Lys Gly Ile Leu Arg Gln Ile Thr Val Asn Asp Leu Pro Val Gly  
 130 135 140

Arg Ser Val Asp Glu Thr Leu Arg Leu Val Gln Ala Phe Gln Tyr Val  
 145 150 155 160

Asp Lys His Gly Glu Val Cys Pro Ala Gly Trp Thr Pro Gly Lys Ala  
165 170 175

Thr Ile Lys Pro Gly Val Lys Asp Ser Lys Glu Tyr Phe Ser Lys Ala  
180 185 190

Asn

<210> 7  
<211> 693  
<212> DNA  
<213> *Ostertagia ostertagi*

<220>  
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<222> (1)..(693)

<220>  
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<222> (11)..(11)  
<223> n is a, c, g, or t

<220>  
<221> misc\_feature  
<222> (35)..(36)  
<223> n is a, c, g, or t

<220>  
<221> misc\_feature  
<222> (41)..(41)  
<223> n is a, c, g, or t

<220>  
<221> misc\_feature  
<222> (43)..(43)  
<223> n is a, c, g, or t

<220>  
<221> misc\_feature  
<222> (53)..(53)  
<223> n is a, c, g, or t

<220>  
<221> misc\_feature  
<222> (67)..(67)  
<223> n is a, c, g, or t

<220>  
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<222> (199)..(199)  
<223> n is a, c, g, or t

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<220>
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<222> (555)..(555)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (590)..(590)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (648)..(648)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (662)..(662)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (669)..(669)
<223> n is a, c, g, or t

<220>
<221> misc_feature
<222> (682)..(682)
<223> n is a, c, g, or t

<400> 7
cta act cct tng cat cca acg cgt tgg gag ctc tnn cta tng ngg gaa 48
Leu Thr Pro Xaa His Pro Thr Arg Trp Glu Leu Xaa Leu Xaa Xaa Glu
1 5 10 15
ttt cna tgt ggt ggc gac nac tcc tgg agc ccg tca gta tcg gcg gaa 96
Leu Xaa Cys Gly Gly Asp Xaa Ser Trp Ser Pro Ser Val Ser Ala Glu
20 25 30
ttc gcg gcc gcg tcg acc gtg ggt gtg gcc ctc gcg gtc cac caa aca 144
Phe Ala Ala Ala Ser Thr Val Gly Val Ala Leu Ala Val His Gln Thr
35 40 45
ctt gac ctg ctt cct ctg aag cca cgc aag gag tac gtc ttc cgc ttt 192
Leu Asp Leu Leu Pro Leu Lys Pro Arg Lys Glu Tyr Val Phe Arg Phe
50 55 60
gaa gga nat gtt cac tcc gga atc ccg ctc cca acc gac acc acc atc 240
Glu Gly Xaa Val His Ser Gly Ile Pro Leu Pro Thr Asp Thr Thr Ile
65 70 75 80
tct cgc ata cag gct atg gta cat gtc cag atc cct gac gac cac cac 288
Ser Arg Ile Gln Ala Met Val His Val Gln Ile Pro Asp Asp His His
85 90 95
gcc att ctc aag ctg aga gat gtt cgc ttt gct act gga gaa gac gaa 336
Ala Ile Leu Lys Leu Arg Asp Val Arg Phe Ala Thr Gly Glu Asp Glu
100 105 110

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cgc aga gaa ctc ttc aaa ccg atc gat gac ctg aaa atg cgc aca atc	384
Arg Arg Glu Leu Phe Lys Pro Ile Asp Asp Leu Lys Met Arg Thr Ile	
115 120 125	
tca agg gag cac ctc gat ctc ctt gag ttg cca gtc cgt ttt gtc tac	432
Ser Arg Glu His Leu Asp Leu Leu Glu Leu Pro Val Arg Phe Val Tyr	
130 135 140	
aag aac ggc atg att tcc gat gta atc ttt gtc gac aag gag gag acc	480
Lys Asn Gly Met Ile Ser Asp Val Ile Phe Val Asp Lys Glu Glu Thr	
145 150 155 160	
tgg tcc cgc cag cgt gaa gcc gat ctg tca tca aca tgc tcc act tta	528
Trp Ser Arg Gln Arg Glu Ala Asp Leu Ser Ser Thr Cys Ser Thr Leu	
165 170 175	
acc tcc aca aga tgg gac gaa ctg acn agc ttt aca atg gac agg tcc	576
Thr Ser Thr Arg Trp Asp Glu Leu Thr Ser Phe Thr Met Asp Arg Ser	
180 185 190	
aag gtg gac ccg tng aca aac gag tac ttt cac tgg tta ccc gaa ccg	624
Lys Val Asp Pro Xaa Thr Asn Glu Tyr Phe His Trp Leu Pro Glu Pro	
195 200 205	
aac cca ttc gaa ggg aaa ctt gtn aag gtt ggc tta cnc cgg ttn tta	672
Asn Pro Phe Glu Gly Lys Leu Val Lys Val Gly Leu Xaa Arg Xaa Leu	
210 215 220	
aag aaa aaa ngg acc ttt tgg	693
Lys Lys Lys Xaa Thr Phe Trp	
225 230	

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<210> 8
<211> 231
<212> PRT
<213> Ostertagia ostertagia

<220>
<221> misc_feature
<222> (4)..(4)
<223> The 'Xaa' at location 4 stands for Trp, Ser, or Leu.

<220>
<221> misc_feature
<222> (12)..(12)
<223> The 'Xaa' at location 12 stands for Tyr, Trp, Cys, Ser, Leu, or
Phe.

<220>
<221> misc_feature
<222> (14)..(14)
<223> The 'Xaa' at location 14 stands for Trp, Ser, or Leu.

<220>
<221> misc_feature
<222> (15)..(15)
<223> The 'Xaa' at location 15 stands for Arg, Gly, or Trp.

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<220>  
<221> misc\_feature  
<222> (18)..(18)  
<223> The 'Xaa' at location 18 stands for Gln, Arg, Pro, or Leu.

<220>  
<221> misc\_feature  
<222> (23)..(23)  
<223> The 'Xaa' at location 23 stands for Asn, Asp, His, or Tyr.

<220>  
<221> misc\_feature  
<222> (67)..(67)  
<223> The 'Xaa' at location 67 stands for Asn, Asp, His, or Tyr.

<220>  
<221> misc\_feature  
<222> (197)..(197)  
<223> The 'Xaa' at location 197 stands for Trp, Ser, or Leu.

<220>  
<221> misc\_feature  
<222> (221)..(221)  
<223> The 'Xaa' at location 221 stands for His, Arg, Pro, or Leu.

<220>  
<221> misc\_feature  
<222> (223)..(223)  
<223> The 'Xaa' at location 223 stands for Leu, or Phe.

<220>  
<221> misc\_feature  
<222> (228)..(228)  
<223> The 'Xaa' at location 228 stands for Arg, Gly, or Trp.

&lt;400&gt; 8

Leu Thr Pro Xaa His Pro Thr Arg Trp Glu Leu Xaa Leu Xaa Xaa Glu  
1 5 10 15

Leu Xaa Cys Gly Gly Asp Xaa Ser Trp Ser Pro Ser Val Ser Ala Glu  
20 25 30

Phe Ala Ala Ala Ser Thr Val Gly Val Ala Leu Ala Val His Gln Thr  
35 40 45

Leu Asp Leu Leu Pro Leu Lys Pro Arg Lys Glu Tyr Val Phe Arg Phe  
50 55 60

Glu Gly Xaa Val His Ser Gly Ile Pro Leu Pro Thr Asp Thr Thr Ile  
65 70 75 80

Ser Arg Ile Gln Ala Met Val His Val Gln Ile Pro Asp Asp His His  
85 90 95

Ala Ile Leu Lys Leu Arg Asp Val Arg Phe Ala Thr Gly Glu Asp Glu  
 100 105 110

Arg Arg Glu Leu Phe Lys Pro Ile Asp Asp Leu Lys Met Arg Thr Ile  
 115 120 125

Ser Arg Glu His Leu Asp Leu Leu Glu Leu Pro Val Arg Phe Val Tyr  
 130 135 140

Lys Asn Gly Met Ile Ser Asp Val Ile Phe Val Asp Lys Glu Glu Thr  
 145 150 155 160

Trp Ser Arg Gln Arg Glu Ala Asp Leu Ser Ser Thr Cys Ser Thr Leu  
 165 170 175

Thr Ser Thr Arg Trp Asp Glu Leu Thr Ser Phe Thr Met Asp Arg Ser  
 180 185 190

Lys Val Asp Pro Xaa Thr Asn Glu Tyr Phe His Trp Leu Pro Glu Pro  
 195 200 205

Asn Pro Phe Glu Gly Lys Leu Val Lys Val Gly Leu Xaa Arg Xaa Leu  
 210 215 220

Lys Lys Lys Xaa Thr Phe Trp  
 225 230

<210> 9  
 <211> 763  
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 <213> *Ostertagia ostertagii*

<220>  
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 <222> (11) .. (706)

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 Met Ser Ala Ala Val Val Val Ala Val Leu Leu Ala Leu  
 1 5 10

ttc tcc tat gcc gaa gca ggc ttt tgt tgt ccg aat agt cta agc caa 97  
 Phe Ser Tyr Ala Glu Ala Gly Phe Cys Cys Pro Asn Ser Leu Ser Gln  
 15 20 25

agt gac agc gcg agg cag att ttc ctc gat ttt cac aat gat gtt cgt 145  
 Ser Asp Ser Ala Arg Gln Ile Phe Leu Asp Phe His Asn Asp Val Arg  
 30 35 40 45

cga aat ata gca ctt gga aat ggt ttg ata aac tgg aca gta aat gca	193
Arg Asn Ile Ala Leu Gly Asn Gly Leu Ile Asn Trp Thr Val Asn Ala	
50 55 60	
gac gcg gtc att ctt ggt cca gct cag aac atg tac aaa gtt gac tgg	241
Asp Ala Val Ile Leu Gly Pro Ala Gln Asn Met Tyr Lys Val Asp Trp	
65 70 75	
gat tgc aac ttg gaa gaa gta gca gca caa cag att gcg cca tgc aat	289
Asp Cys Asn Leu Glu Glu Val Ala Ala Gln Gln Ile Ala Pro Cys Asn	
80 85 90	
gat ccc cta ccg ata aat acc agc ctg gct caa aat atc gct aga tgg	337
Asp Pro Leu Pro Ile Asn Thr Ser Leu Ala Gln Asn Ile Ala Arg Trp	
95 100 105	
ctg tac ttc aaa gac agt gaa gaa gag aca gtt ctg caa caa gta tcg	385
Leu Tyr Phe Lys Asp Ser Glu Glu Glu Thr Val Leu Gln Gln Val Ser	
110 115 120 125	
tgg tat tgg gtc agc gca tcg ctg gga ttt atg aaa ggc acg aaa ctt	433
Trp Tyr Trp Val Ser Ala Ser Leu Gly Phe Met Lys Gly Thr Lys Leu	
130 135 140	
gac caa ttt gct aac cag tgg gct gaa cct cta gca aac att gca aac	481
Asp Gln Phe Ala Asn Gln Trp Ala Glu Pro Leu Ala Asn Ile Ala Asn	
145 150 155	
tat aga aac cga aag gtt gga tgt gcc cat aag atc tgc ccc gct cag	529
Tyr Arg Asn Arg Lys Val Gly Cys Ala His Lys Ile Cys Pro Ala Gln	
160 165 170	
caa aac atg gta gta tcc tgc gtg tat gga agc ccc aaa ctt gca ccg	577
Gln Asn Met Val Val Ser Cys Val Tyr Gly Ser Pro Lys Leu Ala Pro	
175 180 185	
aac gaa gtt atc tgg cag gaa gga aag gct tgt gtg tgc gac gct cgt	625
Asn Glu Val Ile Trp Gln Glu Gly Lys Ala Cys Val Cys Asp Ala Arg	
190 195 200 205	
cca gat tca ttc tgc tgc gac aac ctg tgt gac acg cga gat gct gcg	673
Pro Asp Ser Phe Cys Cys Asp Asn Leu Cys Asp Thr Arg Asp Ala Ala	
210 215 220	
agt gtt cgc cac cag tgt tgc gcg tgc cca tga agcgaaaaga aattggtagt	726
Ser Val Arg His Gln Cys Cys Ala Ser Pro	
225 230	
caccccgaaat aaaatattca tgcaaaaaaaa aaaaaaaaa	763

<210> 10  
 <211> 231  
 <212> PRT  
 <213> Ostertagia ostertagi

<400> 10

Met Ser Ala Ala Val Val Val Ala Val Leu Leu Ala Leu Phe Ser Tyr  
 1 5 10 15

Ala Glu Ala Gly Phe Cys Cys Pro Asn Ser Leu Ser Gln Ser Asp Ser  
20 25 30

Ala Arg Gln Ile Phe Leu Asp Phe His Asn Asp Val Arg Arg Asn Ile  
35 40 45

Ala Leu Gly Asn Gly Leu Ile Asn Trp Thr Val Asn Ala Asp Ala Val  
50 55 60

Ile Leu Gly Pro Ala Gln Asn Met Tyr Lys Val Asp Trp Asp Cys Asn  
65 70 75 80

Leu Glu Glu Val Ala Ala Gln Gln Ile Ala Pro Cys Asn Asp Pro Leu  
85 90 95

Pro Ile Asn Thr Ser Leu Ala Gln Asn Ile Ala Arg Trp Leu Tyr Phe  
100 105 110

Lys Asp Ser Glu Glu Glu Thr Val Leu Gln Gln Val Ser Trp Tyr Trp  
115 120 125

Val Ser Ala Ser Leu Gly Phe Met Lys Gly Thr Lys Leu Asp Gln Phe  
130 135 140

Ala Asn Gln Trp Ala Glu Pro Leu Ala Asn Ile Ala Asn Tyr Arg Asn  
145 150 155 160

Arg Lys Val Gly Cys Ala His Lys Ile Cys Pro Ala Gln Gln Asn Met  
165 170 175

Val Val Ser Cys Val Tyr Gly Ser Pro Lys Leu Ala Pro Asn Glu Val  
180 185 190

Ile Trp Gln Glu Gly Lys Ala Cys Val Cys Asp Ala Arg Pro Asp Ser  
195 200 205

Phe Cys Cys Asp Asn Leu Cys Asp Thr Arg Asp Ala Ala Ser Val Arg  
210 215 220

His Gln Cys Cys Ala Ser Pro  
225 230

<210> 11  
<211> 893  
<212> DNA  
<213> *Ostertagia ostertagi*

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<222> (1) .. (684

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<223> n is a, c, q, or t
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<220>
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<222> (858)..(858)
<223> n is a, c, g, or t
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Met Lys Leu Val Val Leu Cys Val Leu Cys Gly Ile Ala Leu Ala Ala  
1 5 10 15

ccg aga cag aaa cgc ctt act gtg ggc acg atc gct gtc acc gga gga 96  
 Pro Arg Gln Lys Arg Leu Thr Val Gly Thr Ile Ala Val Thr Gly Gly  
 20 25 30

gtc ggc gga tcc acg ggg tgt gta gtg act gga aat gtc ctc tac gca 144  
 Val Gly Gly Ser Thr Gly Cys Val Val Thr Gly Asn Val Leu Tyr Ala  
 35 40 45

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aac ggt ttc cgc ctt cgt gaa ctc aac cca tcg gag cag caa gaa ctc 192
Asn Gly Phe Arg Leu Arg Glu Leu Asn Pro Ser Glu Gln Gln Glu Leu
      50           55           60

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gta aac tat gag aag cag gtg gcc gac tac aaa gcg gct gtg aag caa 240  
 Val Asn Tyr Glu Lys Gln Val Ala Asp Tyr Lys Ala Ala Val Lys Gln  
 65 70 75 80

gcc ctc aag gaa cgc cag gaa agc ctg aaa tcg cgc atg gct ggt aag 288  
Ala Leu Lys Glu Arg Gln Glu Ser Leu Lys Ser Arg Met Ala Gly Lys  
85 90 95

aag gag aag got gtg act ccc aag gag gaa gat cta ccc aaa gct cca 336  
Lys Glu Lys Ala Val Thr Pro Lys Glu Glu Asp Leu Pro Lys Ala Pro  
100 105 110

cag aag ccc tca ttc tgc act gag gac gac acc acc cag ttc tac ttt 384  
 Gln Lys Pro Ser Phe Cys Thr Glu Asp Asp Thr Thr Gln Phe Tyr Phe  
 115 120 125

gat gga tgc atg gtt cag ggc aac aag gtc tac gtt ggc aac aca ttc 432  
 Asp Gly Cys Met Val Gln Gly Asn Lys Val Tyr Val Gly Asn Thr Phe  
 130 135 140

gcg cgc gat ttg gac cag aac gag att caa gag ctg aag gag ttt gag 480  
 Ala Arg Asp Leu Asp Gln Asn Glu Ile Gln Glu Leu Lys Glu Phe Glu  
 145 150 155 160

aag aag cag act gtc tac cag gaa tac gtc cag aag cag att caa gcg Lys Lys Gln Thr Val Tyr Gln Glu Tyr Val Gln Lys Gln Ile Gln Ala 165 170 175	528
caa gtg agc aat ctg ttc ggc ggt gcc gac ttc ttt tca tcg ttc ttc Gln Val Ser Asn Leu Phe Gly Gly Ala Asp Phe Phe Ser Ser Phe Phe 180 185 190	576
aac ggc gga tct gag aaa ggc tct tca acc acc act gtg gcc cca gtg Asn Gly Gly Ser Glu Lys Gly Ser Ser Thr Thr Val Ala Pro Val 195 200 205	624
ctt cct gaa gat gca cca gaa caa cca gct ggg ccc aac ttt tgc aca Leu Pro Glu Asp Ala Pro Glu Gln Pro Ala Gly Pro Asn Phe Cys Thr 210 215 220	672
agg atc tat tga tgggtattt ttatgatgac aaagtattta aataaatgca Arg Ile Tyr 225	724
gtagttgcct gttgctgtga attccacagc actcctactc acgggtgtcga ctggtgattt agtcacttta tttgcaatat ttttatgng ttacccgcaat tcgttgtata tttgtgttat aaacatttaac atcnaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa	784 844 893
<210> 12 <211> 227 <212> PRT <213> <i>Ostertagia ostertagi</i>	
<400> 12	
Met Lys Leu Val Val Leu Cys Val Leu Cys Gly Ile Ala Leu Ala Ala 1 5 10 15	
Pro Arg Gln Lys Arg Leu Thr Val Gly Thr Ile Ala Val Thr Gly Gly 20 25 30	
Val Gly Gly Ser Thr Gly Cys Val Val Thr Gly Asn Val Leu Tyr Ala 35 40 45	
Asn Gly Phe Arg Leu Arg Glu Leu Asn Pro Ser Glu Gln Gln Glu Leu 50 55 60	
Val Asn Tyr Glu Lys Gln Val Ala Asp Tyr Lys Ala Ala Val Lys Gln 65 70 75 80	
Ala Leu Lys Glu Arg Gln Glu Ser Leu Lys Ser Arg Met Ala Gly Lys 85 90 95	

Lys Glu Lys Ala Val Thr Pro Lys Glu Glu Asp Leu Pro Lys Ala Pro  
 100 105 110

Gln Lys Pro Ser Phe Cys Thr Glu Asp Asp Thr Thr Gln Phe Tyr Phe  
 115 120 125

Asp Gly Cys Met Val Gln Gly Asn Lys Val Tyr Val Gly Asn Thr Phe  
 130 135 140

Ala Arg Asp Leu Asp Gln Asn Glu Ile Gln Glu Leu Lys Glu Phe Glu  
 145 150 155 160

Lys Lys Gln Thr Val Tyr Gln Glu Tyr Val Gln Lys Gln Ile Gln Ala  
 165 170 175

Gln Val Ser Asn Leu Phe Gly Gly Ala Asp Phe Phe Ser Ser Phe Phe  
 180 185 190

Asn Gly Gly Ser Glu Lys Gly Ser Ser Thr Thr Thr Val Ala Pro Val  
 195 200 205

Leu Pro Glu Asp Ala Pro Glu Gln Pro Ala Gly Pro Asn Phe Cys Thr  
 210 215 220

Arg Ile Tyr  
 225

<210> 13  
 <211> 1761  
 <212> DNA  
 <213> *Ostertagia ostertagi*

<220>  
 <221> CDS  
 <222> (1)..(1725)

<400> 13  
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 Met Arg Leu Ile Leu Leu Ile Leu Leu Val Val Ala Thr Asn Gly  
 1 5 10 15  
 ggc ata att gac aaa ctg aaa gga ttg ttc act gga gaa ggc ggc ttt 96  
 Gly Ile Ile Asp Lys Leu Lys Gly Leu Phe Thr Gly Glu Gly Phe  
 20 25 30  
 gga caa aaa gtg aag aat gca act gct gtt ggc ttc aaa aag ctc ttc 144  
 Gly Gln Lys Val Lys Asn Ala Thr Ala Val Gly Phe Lys Lys Leu Phe  
 35 40 45

gaa aac acg gca ctc ttc aga atc aat gat aag atc agg agc atg aag Glu Asn Thr Ala Leu Phe Arg Ile Asn Asp Lys Ile Arg Ser Met Lys 50 55 60	192
gaa aaa gtg ttg aag acc ttg gaa cta tca cca gca atg atg aag tca Glu Lys Val Leu Lys Thr Leu Glu Leu Ser Pro Ala Met Met Lys Ser 65 70 75 80	240
ctg caa kmg agg cta rwd aaw tsg cgr cck yct rma grw cga yma wrt Leu Gln Xaa Arg Leu Xaa Xaa Xaa Arg Xaa Xaa Xaa Arg Xaa Xaa 85 90 95	288
rsr mga gmt sss aga crc gtw kka ygc rag gtc art aaa aat agt gag Xaa Xaa Xaa Xaa Arg Xaa Xaa Xaa Val Xaa Lys Asn Ser Glu 100 105 110	336
gtt gac caa tac ctc tac caa ggc gac atg gtt tta aca gag gag caa Val Asp Gln Tyr Leu Tyr Gln Gly Asp Met Val Leu Thr Glu Glu Gln 115 120 125	384
gcc gat gag atc gtt gag gac ata gaa gat cag gtc gcc ggt gga aat Ala Asp Glu Ile Val Glu Asp Ile Glu Asp Gln Val Ala Gly Gly Asn 130 135 140	432
cgt aca aaa cgt caa gca ttc aag gat cat aaa tat ccc aaa acg ttg Arg Thr Lys Arg Gln Ala Phe Lys Asp His Lys Tyr Pro Lys Thr Leu 145 150 155 160	480
tgg tca caa gga gtc aac tac tac ttc cat gat atg gcc agt aag cag Trp Ser Gln Gly Val Asn Tyr Tyr Phe His Asp Met Ala Ser Lys Gln 165 170 175	528
atg aaa agc gta ttc gta aaa gga gcg aaa tgg tgg gaa aag gac acg Met Lys Ser Val Phe Val Lys Gly Ala Lys Trp Trp Glu Lys Asp Thr 180 185 190	576
tgt atc aat ttc acg gag aac cgt tct gcc gaa gac cga att atg gta Cys Ile Asn Phe Thr Glu Asn Arg Ser Ala Glu Asp Arg Ile Met Val 195 200 205	624
ttc cca cag aaa gga tgt tgg tca aat ata gga aaa atc ggt ggc gaa Phe Pro Gln Lys Gly Cys Trp Ser Asn Ile Gly Lys Ile Gly Gly Glu 210 215 220	672
caa aag att tcg ttg gga gga ggt tgt cat tcg gta tcc att gct gcg Gln Lys Ile Ser Leu Gly Gly Cys His Ser Val Ser Ile Ala Ala 225 230 235 240	720
cat gag atc ggt cac gca att gga ttc ttc cat act atg tcc cgt cac His Glu Ile Gly His Ala Ile Gly Phe Phe His Thr Met Ser Arg His 245 250 255	768
gat cgc gat gaa ttc atc acc gta aac atg cac aat gtt gat gta cac Asp Arg Asp Glu Phe Ile Thr Val Asn Met His Asn Val Asp Val His 260 265 270	816
tgg ctg agt caa ttt aat aaa gaa acg acg aag aga aat gat aat tat Trp Ile Ser Gln Phe Asn Lys Glu Thr Thr Lys Arg Asn Asp Asn Tyr 275 280 285	864

gga atg acg tac gac tac ggt agc att atg cat tac ggt gga acc agt Gly Met Thr Tyr Asp Tyr Gly Ser Ile Met His Tyr Gly Gly Thr Ser 290 295 300	912
gca tcg tac aat aat aag cca aca atg gtg ccg ttt gat gtg gac tat Ala Ser Tyr Asn Asn Lys Pro Thr Met Val Pro Phe Asp Val Asp Tyr 305 310 315 320	960
cag caa acc ctt ggc tct cca ttc att tct ttc att gaa ctt tcc atg Gln Gln Thr Leu Gly Ser Pro Phe Ile Ser Phe Ile Glu Leu Ser Met 325 330 335	1008
att aat gaa cac tac aaa tgc aaa gag aac tgc aat cca gct aag tcg Ile Asn Glu His Tyr Lys Cys Lys Glu Asn Cys Asn Pro Ala Lys Ser 340 345 350	1056
gct aaa tgc gaa atg ggc gga ttc cct cat ccc cga gac tgc agc aaa Ala Lys Cys Glu Met Gly Gly Phe Pro His Pro Arg Asp Cys Ser Lys 355 360 365	1104
tgt atc tgt cct ggt gga tac gcc gga gct cga tgc acc gaa aga cca Cys Ile Cys Pro Gly Gly Tyr Ala Gly Ala Arg Cys Thr Glu Arg Pro 370 375 380	1152
tca ggg tgt ggc agt gca att caa gct tcg tcc gat tgg aag acc tta Ser Gly Cys Gly Ser Ala Ile Gln Ala Ser Ser Asp Trp Lys Thr Leu 385 390 395 400	1200
caa gat acc ctt ggc aag gat gat gat gaa gaa cga gag gac ttc gag Gln Asp Thr Leu Gly Lys Asp Asp Asp Glu Glu Arg Glu Asp Phe Glu 405 410 415	1248
aca tgt aat tac tgg att gaa tct cct gcc gga acm gaa atc gaa gtt Thr Cys Asn Tyr Trp Ile Glu Ser Pro Ala Gly Xaa Glu Ile Glu Val 420 425 430	1296
agg tta ttg gat ttc acg agg ggt gtt tct gtc gat gga tgc aaa ttt Arg Leu Leu Asp Phe Thr Arg Gly Val Ser Val Asp Gly Cys Lys Phe 435 440 445	1344
gcc ggt gta gag atc aag acc aat aag gat caa aca ctc act ggc tac Ala Gly Val Glu Ile Lys Thr Asn Lys Asp Gln Thr Leu Thr Gly Tyr 450 455 460	1392
agg ttc tgc aca gct ggc gca gct ggc ata gca ctt cgt tct tac acg Arg Phe Cys Thr Ala Gly Ala Ala Gly Ile Ala Leu Arg Ser Tyr Thr 465 470 475 480	1440
aat cgc gtc cca ata atg aca tac aac aga ttt ggt caa tcg acg act Asn Arg Val Pro Ile Met Thr Tyr Asn Arg Phe Gly Gln Ser Thr Thr 485 490 495	1488
gtt ctc gaa tac cga cac gtt ccg gcg agt ggc cca aga acg ccc tca Val Leu Glu Tyr Arg His Val Pro Ala Ser Ala Pro Arg Thr Pro Ser 500 505 510	1536
cct cca tct gct aca act cgt gct tct att act act act act act acg Pro Pro Ser Ala Thr Thr Arg Ala Ser Ile Thr Thr Thr Thr Thr 515 520 525	1584

aag aaa ccc agc tct act gct gcc ttt aaa tgc gag gac aac cac act 1632  
Lys Lys Pro Ser Ser Thr Ala Ala Phe Lys Cys Glu Asp Asn His Thr  
530 535 540

tgt ccc tca ctt gta gcg agc ggt ttc tgc aaa ggg cca ctc tca gag 1680  
Cys Pro Ser Leu Val Ala Ser Gly Phe Cys Lys Gly Pro Leu Ser Glu  
545 550 555 560

gct acc aag aag aaa gtg tgt cca aag tcg tgt gga ctc tgc tga 1725  
Ala Thr Lys Lys Val Cys Pro Lys Ser Cys Gly Leu Cys  
565 570

tacaacactt tctctgtaat aaaatctgaa caattc 1761

<210> 14  
<211> 574  
<212> PRT  
<213> Ostertagia ostertagii

<220>  
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<222> (83)..(83)  
<223> The 'Xaa' at location 83 stands for Glu, Ala, or Ser.

<220>  
<221> misc\_feature  
<222> (86)..(86)  
<223> The 'Xaa' at location 86 stands for Glu, Val, Lys, or Met.

<220>  
<221> misc\_feature  
<222> (87)..(87)  
<223> The 'Xaa' at location 87 stands for Lys, or Asn.

<220>  
<221> misc\_feature  
<222> (88)..(88)  
<223> The 'Xaa' at location 88 stands for Trp, or Ser.

<220>  
<221> misc\_feature  
<222> (90)..(90)  
<223> The 'Xaa' at location 90 stands for Pro.

<220>  
<221> misc\_feature  
<222> (91)..(91)  
<223> The 'Xaa' at location 91 stands for Pro, or Ser.

<220>  
<221> misc\_feature  
<222> (92)..(92)  
<223> The 'Xaa' at location 92 stands for Glu, Ala, Lys, or Thr.

<220>  
<221> misc\_feature  
<222> (93)..(93)  
<223> The 'Xaa' at location 93 stands for Gly, Glu, or Asp.

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<220>
<221> misc_feature
<222> (95)..(95)
<223> The 'Xaa' at location 95 stands for Gln, Pro, or Ser.

<220>
<221> misc_feature
<222> (96)..(96)
<223> The 'Xaa' at location 96 stands for Ser, Asn, Cys, or Tyr.

<220>
<221> misc_feature
<222> (97)..(97)
<223> The 'Xaa' at location 97 stands for Gly, Ala, Arg, or Thr.

<220>
<221> misc_feature
<222> (98)..(98)
<223> The 'Xaa' at location 98 stands for Arg.

<220>
<221> misc_feature
<222> (99)..(99)
<223> The 'Xaa' at location 99 stands for Asp, or Ala.

<220>
<221> misc_feature
<222> (100)..(100)
<223> The 'Xaa' at location 100 stands for Gly, Ala, Arg, or Pro.

<220>
<221> misc_feature
<222> (102)..(102)
<223> The 'Xaa' at location 102 stands for Arg, or His.

<220>
<221> misc_feature
<222> (103)..(103)
<223> The 'Xaa' at location 103 stands for Val.

<220>
<221> misc_feature
<222> (104)..(104)
<223> The 'Xaa' at location 104 stands for Gly, Val, or Leu.

<220>
<221> misc_feature
<222> (105)..(105)
<223> The 'Xaa' at location 105 stands for Arg, or Cys.

<220>
<221> misc_feature
<222> (106)..(106)
<223> The 'Xaa' at location 106 stands for Glu, or Lys.

<220>
<221> misc_feature
<222> (108)..(108)
<223> The 'Xaa' at location 108 stands for Ser, or Asn.
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<220>  
<221> misc\_feature  
<222> (428)..(428)  
<223> The 'Xaa' at location 428 stands for Thr.

<400> 14

Met Arg Leu Ile Leu Leu Ile Leu Leu Leu Val Val Ala Thr Asn Gly  
1 5 10 15

Gly Ile Ile Asp Lys Leu Lys Gly Leu Phe Thr Gly Glu Gly Gly Phe  
20 25 30

Gly Gln Lys Val Lys Asn Ala Thr Ala Val Gly Phe Lys Lys Leu Phe  
35 40 45

Glu Asn Thr Ala Leu Phe Arg Ile Asn Asp Lys Ile Arg Ser Met Lys  
50 55 60

Glu Lys Val Leu Lys Thr Leu Glu Leu Ser Pro Ala Met Met Lys Ser  
65 70 75 80

Leu Gln Xaa Arg Leu Xaa Xaa Xaa Arg Xaa Xaa Xaa Xaa Arg Xaa Xaa  
85 90 95

Xaa Xaa Xaa Xaa Arg Xaa Xaa Xaa Xaa Val Xaa Lys Asn Ser Glu  
100 105 110

Val Asp Gln Tyr Leu Tyr Gln Gly Asp Met Val Leu Thr Glu Glu Gln  
115 120 125

Ala Asp Glu Ile Val Glu Asp Ile Glu Asp Gln Val Ala Gly Gly Asn  
130 135 140

Arg Thr Lys Arg Gln Ala Phe Lys Asp His Lys Tyr Pro Lys Thr Leu  
145 150 155 160

Trp Ser Gln Gly Val Asn Tyr Tyr Phe His Asp Met Ala Ser Lys Gln  
165 170 175

Met Lys Ser Val Phe Val Lys Gly Ala Lys Trp Trp Glu Lys Asp Thr  
180 185 190

Cys Ile Asn Phe Thr Glu Asn Arg Ser Ala Glu Asp Arg Ile Met Val  
195 200 205

Phe Pro Gln Lys Gly Cys Trp Ser Asn Ile Gly Lys Ile Gly Gly Glu  
210 215 220

Gln Lys Ile Ser Leu Gly Gly Cys His Ser Val Ser Ile Ala Ala  
225 230 235 240

His Glu Ile Gly His Ala Ile Gly Phe Phe His Thr Met Ser Arg His  
245 250 255

Asp Arg Asp Glu Phe Ile Thr Val Asn Met His Asn Val Asp Val His  
260 265 270

Trp Leu Ser Gln Phe Asn Lys Glu Thr Thr Lys Arg Asn Asp Asn Tyr  
275 280 285

Gly Met Thr Tyr Asp Tyr Gly Ser Ile Met His Tyr Gly Gly Thr Ser  
290 295 300

Ala Ser Tyr Asn Asn Lys Pro Thr Met Val Pro Phe Asp Val Asp Tyr  
305 310 315 320

Gln Gln Thr Leu Gly Ser Pro Phe Ile Ser Phe Ile Glu Leu Ser Met  
325 330 335

Ile Asn Glu His Tyr Lys Cys Lys Glu Asn Cys Asn Pro Ala Lys Ser  
340 345 350

Ala Lys Cys Glu Met Gly Phe Pro His Pro Arg Asp Cys Ser Lys  
355 360 365

Cys Ile Cys Pro Gly Gly Tyr Ala Gly Ala Arg Cys Thr Glu Arg Pro  
370 375 380

Ser Gly Cys Gly Ser Ala Ile Gln Ala Ser Ser Asp Trp Lys Thr Leu  
385 390 395 400

Gln Asp Thr Leu Gly Lys Asp Asp Glu Glu Arg Glu Asp Phe Glu  
405 410 415

Thr Cys Asn Tyr Trp Ile Glu Ser Pro Ala Gly Xaa Glu Ile Glu Val  
420 425 430

Arg Leu Leu Asp Phe Thr Arg Gly Val Ser Val Asp Gly Cys Lys Phe  
435 440 445

Ala Gly Val Glu Ile Lys Thr Asn Lys Asp Gln Thr Leu Thr Gly Tyr  
 450 455 460

Arg Phe Cys Thr Ala Gly Ala Ala Gly Ile Ala Leu Arg Ser Tyr Thr  
 465 470 475 480

Asn Arg Val Pro Ile Met Thr Tyr Asn Arg Phe Gly Gln Ser Thr Thr  
 485 490 495

Val Leu Glu Tyr Arg His Val Pro Ala Ser Ala Pro Arg Thr Pro Ser  
 500 505 510

Pro Pro Ser Ala Thr Thr Arg Ala Ser Ile Thr Thr Thr Thr Thr Thr  
 515 520 525

Lys Lys Pro Ser Ser Thr Ala Ala Phe Lys Cys Glu Asp Asn His Thr  
 530 535 540

Cys Pro Ser Leu Val Ala Ser Gly Phe Cys Lys Gly Pro Leu Ser Glu  
 545 550 555 560

Ala Thr Lys Lys Lys Val Cys Pro Lys Ser Cys Gly Leu Cys  
 565 570

<210> 15  
 <211> 24  
 <212> DNA  
 <213> Artificial

<220>  
 <223> primer: Lambdagt11F

<220>  
 <221> misc\_feature  
 <223> Lambdagt11F

<400> 15  
 ggtggcgacg actcctggag cccg

24

<210> 16  
 <211> 24  
 <212> DNA  
 <213> Artificial

<220>  
 <223> primer: Lambdagt11R

<400> 16  
 ttgacaccag accaactggg aatg

24

<210> 17  
<211> 20  
<212> DNA  
<213> Artificial  
  
<220>  
<223> primer: SP6  
  
<400> 17  
atttaggtga cactatagaa

20

<210> 18  
<211> 22  
<212> DNA  
<213> Artificial  
  
<220>  
<223> primer: T7  
  
<400> 18  
gtaatacgac tcactatagg gc

22

<210> 19  
<211> 21  
<212> DNA  
<213> Artificial  
  
<220>  
<223> primer: 24kForw  
  
<400> 19  
gaattcatga agttggtcgt g

21

<210> 20  
<211> 22  
<212> DNA  
<213> Artificial  
  
<220>  
<223> primer: 24kRev  
  
<400> 20  
ctcgagtcaa tagatccttg tg

22

<210> 21  
<211> 36  
<212> DNA  
<213> Artificial  
  
<220>  
<223> primer: AAP

<220>  
<221> misc\_feature  
<222> (24)..(25)  
<223> n = Inosine

<220>  
<221> misc\_feature  
<222> (29)..(30)  
<223> n = Inosine

<220>  
<221> misc\_feature  
<222> (34)..(35)  
<223> n = Inosine

<400> 21  
ggccacgcgt cgactagtagac gggnnnnnnn gggnnng 36

<210> 22  
<211> 32  
<212> DNA  
<213> Artificial

<220>  
<223> primer: UAP

<400> 22  
cuacuacuac uaggccacgc gtcgactagt ac 32

<210> 23  
<211> 21  
<212> DNA  
<213> Artificial

<220>  
<223> primer: 65Rev1

<400> 23  
cagcaatgga taccgaatga c 21

<210> 24  
<211> 22  
<212> DNA  
<213> Artificial

<220>  
<223> primer: 65Rev2

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